
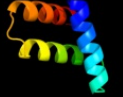
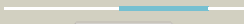












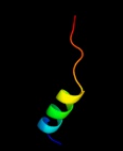


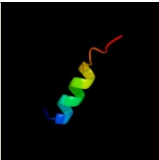
Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c2oxlA_	 Alignment		72.0	19	PDB header: gene regulation Chain: A: PDB Molecule: hypothetical protein ymgb; PDBTitle: structure and function of the e. coli protein ymgb: a protein critical2 for biofilm formation and acid resistance
2	d2bida_	 Alignment		31.9	25	Fold: Toxins' membrane translocation domains Superfamily: Bcl-2 inhibitors of programmed cell death Family: Bcl-2 inhibitors of programmed cell death
3	d1ddba_	 Alignment		30.9	29	Fold: Toxins' membrane translocation domains Superfamily: Bcl-2 inhibitors of programmed cell death Family: Bcl-2 inhibitors of programmed cell death
4	d1r8ja2	 Alignment		8.6	13	Fold: Flavodoxin-like Superfamily: CheY-like Family: N-terminal domain of the circadian clock protein KaiA
5	c2uwjE_	 Alignment		8.5	31	PDB header: chaperone Chain: E: PDB Molecule: type iii export protein psce; PDBTitle: structure of the heterotrimeric complex which regulates2 type iii secretion needle formation
6	c3bbzA_	 Alignment		7.8	53	PDB header: viral protein, replication Chain: A: PDB Molecule: p protein; PDBTitle: structure of the nucleocapsid-binding domain from the mumps2 virus phosphoprotein
7	c3us8A_	 Alignment		7.2	50	PDB header: oxidoreductase Chain: A: PDB Molecule: isocitrate dehydrogenase [nadp]; PDBTitle: crystal structure of an isocitrate dehydrogenase from sinorhizobium2 meliloti 1021
8	c2a40C_	 Alignment		6.3	37	PDB header: structural protein Chain: C: PDB Molecule: wiskott-aldrich syndrome protein family member 2; PDBTitle: ternary complex of the wh2 domain of wave with actin-dnase i

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[c2a40F_](#)

Alignment



6.3

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PDB header:structural protein
Chain: F: **PDB Molecule:**wiskott-aldrich syndrome protein family member 2;
PDBTitle: ternary complex of the wh2 domain of wave with actin-dnase i